- 16. A method for detecting a degree of hybridization between a pr be and a sample comprising a biopolymer, the method comprising
- (a) providing a substrate on which each of a plurality of types of probes is separately immobilized on each different and separate <u>predetermined</u> position, wherein the probes are labeled with a first detectable label;
- (b) providing a sample comprising a biopolymer, wherein the biopolymer is labeled with a second detectable label;
 - (c) contacting the sample with the probe;
- (d) detecting an amount of the probe at each <u>different and separate</u> <u>predetermined</u> position of the substrate by detecting the first detectable label;
- (e) detecting an amount of the sample biopolymer [bound] <u>hybridized</u> to the probe at each <u>different and separate predetermined</u> position of the substrate by detecting the second detectable label; and
- (f) producing a value representing the degree of hybridization between [a] the probe at each different and separate predetermined position and [a] the sample biopolymer by [normalizing] dividing the difference between the amount of the probe detected at each different and separate predetermined position and the amount of the sample biopolymer hybridized to the probe [with] by the amount of the probe.
- 24. A method for detecting [the] a degree of hybridization between an oligonucleotide probe immobilized onto an array and a sample nucleic acid, the method comprising
- (a) providing a substrate on which each of a plurality of types of oligonucleotide probes is separately immobilized on each different and separate predetermined position to form an array, wherein the oligonucleotide probes are labeled with a first detectable label;
- (b) providing a sample comprising a nucleic acid, wherein the nucleic acid is labeled with a second detectable label;
 - (c) contacting the sample with the probe;
- (d) detecting an amount of the probe at each <u>different</u> and <u>separate</u> <u>predetermined</u> position of the substrate by detecting the first detectable label;
- (c) detecting an amount of the sample nucleic acid hybridized to the probe at each <u>different and separate predetermined</u> position of the substrate by detecting the second detectable label; and
- (f) producing a value representing the degree of hybridization between [a] the probe at each different and separate predetermined position and a sample by

[normalizing] dividing the difference between the amount of the probe detected at ach different and separate predetermined position and the amount of the sample nucleic acid hybridized to the probe [with] by the amount of the probe.